Amendments to the Specification:

Please replace paragraph [0176] with the following amended paragraph:

[0176] The human TRAF2 (SEQ ID NO:23) was cloned by PCR from an HL60 cDNA library (for TRAF2 sequence and other details see Rothe et al., 1994; Rothe et al., 1995a; Cheng et al., 1996; Hsu et al., 1996; and Wallach, 1996). The primers used were: a) 30-mer forward primer CAGGATCCTCATGGCTGCAGCTAGCGTGAC (SEQ ID NO:8) corresponding to the coding sequence of hTRAF2 starting from the codon for the first methionine (underlined) and including a linker with BamHI site. b) 32-mer reverse primer GGTCGACTTAGAGCCCTGTCAGGTCCACAATG (SEQ ID NO:9) that includes hTRAF2 gene stop codon (underlined) and a SalI restriction site in its linker. PCR program comprised of an initial denaturation step 2 min. at 94°C followed by 30 cycles of 1 min. at 94°C, 1 min. at 64°C, 1 min. and 40 sec. at 72°C. The amplified human TRAF2 was then inserted into the BamHI-SalI sites of pGBT9 vector in conjunction with GAL 4 DNA Binding domain.